

ERCC

# *Performance of RNA Standards*

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**EXTERNAL RNA CONTROLS CONSORTIUM WORKSHOP**

***NIST***

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## **GOAL:**

**To develop a set RNA standards that will provide a range of signal intensities and/ or gene expression ratios that mimics what is found in “normal” mammalian cells.**

**“Average” cell: Contains 25 pg total RNA  
and 2% or 0.5 pg mRNA**

**#mRNA Molecules/cell: 100,000-500,000**

**Sensitivity Definition: molecules mRNA  
per 100,000 or parts per million**

**Example: 25 ug RNA = 1 million cells or  
>10<sup>11</sup> mRNA molecules. To spike in at  
conc of ~1 part per million use 100,000  
molecules of standard**

## **Testing Performance**

**Spikes: 0, 1, 10, 50, 100.....1000 copies**

**Ratios: 50/1, 25/1, 10/1, 5/1, 2/1, 1/1**

## **Questions:**

**Lowest signal (lowest copy number) detection?**

*Perhaps use absolute numbers only;  $10^4$ ,  $10^5$ ,  $10^6$ , etc.*

**What is dynamic range expected?  $10^0$  to  $10^?$**

**More spikes or ratios required? More the better  
or will more make QC, analysis, etc difficult?**